

# CRF Errors Corrected by the STIC System Branch

Serial Number: 09/033,061

CRF Processing Date: 3/7/98  
 Edited by: AL  
 Verified by: \_\_\_\_\_ (STIC staff)

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☒ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: 21
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☐ Other: \_\_\_\_\_

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/030,061

DATE: 03/07/98  
TIME: 16:28:12

INPUT SET: S2912.raw

This Raw Listing contains the General  
Information Section and those Sequences  
containing ERRORS.

1 SEQUENCE LISTING  
2  
3 (1) General Information:  
4  
5 (i) APPLICANT: GILLISPIE, Matthew Todd  
6 HORWOOD, Nicole Joy  
7 UDAGAWA, Nobuyuki  
8 KURIMOTO, Masashi  
9  
10 (ii) TITLE OF INVENTION: OSTEOCLASTGENIC INHIBITORY AGENT  
11  
12 (iii) NUMBER OF SEQUENCES: 28  
13  
14 (iv) CORRESPONDENCE ADDRESS:  
15 (A) ADDRESSEE: BROWDY AND NEIMARK  
16 (B) STREET: 419 Seventh Street, N.W., Suite 300  
17 (C) CITY: Washington  
18 (D) STATE: D.C.  
19 (E) COUNTRY: USA  
20 (F) ZIP: 20004  
21  
22 (v) COMPUTER READABLE FORM:  
23 (A) MEDIUM TYPE: Floppy disk  
24 (B) COMPUTER: IBM PC compatible  
25 (C) OPERATING SYSTEM: PC-DOS/MS-DOS  
26 (D) SOFTWARE: Patent In Release #1.0, Version #1.30  
27  
28 (vi) CURRENT APPLICATION DATA:  
29 (A) APPLICATION NUMBER:  
30 (B) FILING DATE: 25-FEB-1998  
31  
32 (vii) PRIOR APPLICATION DATA:  
33 (A) APPLICATION NUMBER: JP 55,468/1997  
34 (B) FILING DATE: 25-FEB-1997  
35  
36 (viii) ATTORNEY/AGENT INFORMATION:  
37 (A) NAME: BROWDY, Roger L.  
38 (B) REGISTRATION NUMBER: 25,618  
39 (C) REFERENCE/DOCKET NUMBER: GILLISPIE=1  
40  
41 (ix) TELECOMMUNICATION INFORMATION:  
42 (A) TELEPHONE: (202) 628-5197  
43 (B) TELEFAX: (202) 737-3528  
44

ERRORED SEQUENCES FOLLOW:

RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/030,061DATE: 03/07/98  
TIME: 16:28:14

INPUT SET: S2912.raw

808 (2) INFORMATION FOR SEQ ID NO: 21:

809

810 (i) SEQUENCE CHARACTERISTICS:

811 (A) LENGTH: 157 amino acids

812 (B) TYPE: amino acid

813 (D) TOPOLOGY: linear

814

815 (ii) MOLECULE TYPE: peptide

816

--&gt; 817 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:

818

819 Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn

820 1 5 10 15

821 Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp

822 20 25 30

823 Met Thr Asp Ser Asp Ser Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile

824 35 40 45

825 Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile

826 50 55 60

827 Ser Val Lys Ser Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile

828 65 70 75 80

829 Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys

830 85 90 95

831 Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys

832 100 105 110

833 Met Gln Phe Glu Ser Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu

834 115 120 125

835 Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Leu

836 130 135 140

837 Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp

838 145 150 155

839

PAGE: 1

**SEQUENCE VERIFICATION REPORT**  
**PATENT APPLICATION US/09/030,061**

DATE: 03/07/98  
TIME: 16:28:16

*INPUT SET: S2912.raw*

Line	Error	Original Text
817	Wrong Sequence Number	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 25: